

Substitute SEQUENCE LISTING

<110> Kwon, Byoung

<120> NEW RECEPTOR AND RELATED PRODUCTS AND
METHODS

<130> 740.013US2

<140> 08/955,572

<141> 1997-10-22

<150> 08/461,652

<151> 1995-06-05

<150> 08/122,796

<151> 1993-09-03

<160> 12

<170> FastSEQ for Windows Version 3.0

<210> 1

<211> 838

<212> DNA

<213> Homo sapiens

<400> 1

aatcagcttt	gctagtagcc	tacctgtgcc	agatttcac	atgggaaaca	gctgttaca	60
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ttgtagtaac	tgcccagctg	gtacattctg	tgataataac	aggaatcaga	tttgcagtcc	180
ctgtcctcca	aatagtttct	ccagcgcagg	tgacaaaagg	acctgtgaca	tatgcaggca	240
gtgtaaaggt	gttttcagga	ccaggaagga	gtgttcctcc	accagcaatg	cagagtgtga	300
ctgcactcca	gggtttcact	gcctgggggc	aggatgcagc	atgtgtgaac	aggattgtaa	360
acaagggtcaa	gaactgacaa	aaaaagggtt	taaagactgt	tgctttggga	catttaacga	420
tcagaaacgt	ggcatctgtc	gaccctggac	aaactgttct	ttggatggaa	agtctgtgct	480
tgtgaatggg	acgaaggaga	gggacgtggg	ctgtggacca	tctccagctg	acctctctcc	540
gggagcatcc	tctgtgaccc	cgcctgcccc	tgcgagagag	ccaggacact	ctccgcagat	600
catctccttc	tttcttgccg	tgacgtcgac	tgcggttgct	ttcctgctgt	tcttcctcac	660
gctccgtttc	tctgttggtt	aacggggcag	aaagaaactc	ctgtatatat	tcaaacaacc	720
atztatgaga	ccagtagaaa	ctactcaaga	ggaagatggc	tgtagctgcc	gatttccaga	780
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<210> 2

<211> 255

<212> PRT

<213> Homo sapiens

<400> 2

Met	Gly	Asn	Ser	Cys	Tyr	Asn	Ile	Val	Ala	Thr	Leu	Leu	Leu	Val	Leu
1				5				10						15	
Asn	Phe	Glu	Arg	Thr	Arg	Ser	Leu	Gln	Asp	Pro	Cys	Ser	Asn	Cys	Pro
			20				25						30		
Ala	Gly	Thr	Phe	Cys	Asp	Asn	Asn	Arg	Asn	Gln	Ile	Cys	Ser	Pro	Cys
		35				40					45				
Pro	Pro	Asn	Ser	Phe	Ser	Ser	Ala	Gly	Gly	Gln	Arg	Thr	Cys	Asp	Ile
		50			55					60					
Cys	Arg	Gln	Cys	Lys	Gly	Val	Phe	Arg	Thr	Arg	Lys	Glu	Cys	Ser	Ser
65				70				75				80			
Thr	Ser	Asn	Ala	Glu	Cys	Asp	Cys	Thr	Pro	Gly	Phe	His	Cys	Leu	Gly

<210> 8
<211> 30
<212> DNA
<213> Homo sapiens

<400> 8
ttaagatctc tgcggagagt gtcctggctc

30

<210> 9
<211> 2350
<212> DNA
<213> Mus musculus

<220>
<221> unsure
<222> (1253)...(1255)
<223> (a or g or c or t/u)

<400> 9

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ctccagcata	ggtggacagc	cgaactgtaa	catctgcaga	gtgtgtgcag	gctatttcag	360
gttcaagaag	ttttgtcctc	ctaccacaaa	cgcggagtgt	gagtgcattg	aaggattcca	420
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ggagaaggac	gtggtgtgtg	gacccccctg	ggtgagcttc	tctcccagta	ccaccatttc	660
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tggcgccaag	ataaaacaac	caaaagcctt	gactccggta	ctaattctcc	ctgccggccc	2040
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<210> 10

<211> 256
 <212> PRT
 <213> Mus musculus

<400> 10

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Met Gly Asn Asn Cys Tyr Asn Val Val Val Ile Val Leu Leu Leu Val
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 20      25      30
Pro Gly Thr Phe Cys Arg Lys Tyr Asn Pro Val Cys Lys Ser Cys Pro
 35      40      45
Pro Ser Thr Phe Ser Ser Ile Gly Gly Gln Pro Asn Cys Asn Ile Cys
 50      55      60
Arg Val Cys Ala Gly Tyr Phe Arg Phe Lys Lys Phe Cys Ser Ser Thr
 65      70      75      80
His Asn Ala Glu Cys Glu Cys Ile Glu Gly Phe His Cys Leu Gly Pro
 85      90      95
Gln Cys Thr Arg Cys Glu Lys Asp Cys Arg Pro Gly Gln Glu Leu Thr
 100     105     110
Lys Gln Gly Cys Lys Thr Cys Ser Leu Gly Thr Phe Asn Asp Gln Asn
 115     120     125
Gly Thr Gly Val Cys Arg Pro Trp Thr Asn Cys Ser Leu Asp Gly Arg
 130     135     140
Ser Val Leu Lys Thr Gly Thr Thr Glu Lys Asp Val Val Cys Gly Pro
 145     150     155     160
Pro Val Val Ser Phe Ser Pro Ser Thr Thr Ile Ser Val Thr Pro Glu
 165     170     175
Gly Gly Pro Gly Gly His Ser Leu Gln Val Leu Thr Leu Phe Leu Ala
 180     185     190
Leu Thr Ser Ala Leu Leu Leu Ala Leu Ile Phe Ile Thr Leu Leu Phe
 195     200     205
Ser Val Leu Lys Trp Ile Arg Lys Lys Phe Pro His Ile Phe Lys Gln
 210     215     220
Pro Phe Lys Lys Thr Thr Gly Ala Ala Gln Glu Glu Asp Ala Cys Ser
 225     230     235     240
Cys Arg Cys Pro Gln Glu Glu Glu Gly Gly Gly Gly Gly Tyr Glu Leu
 245     250     255

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<210> 11
 <211> 24
 <212> PRT
 <213> Homo sapiens

<220>

<221> ZN_FING

<222> 2...3, 5...13, 15...17, 19...21, 23

<223> Putative zinc finger structure

<400> 11

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Cys Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
 1      5      10      15
Xaa His Xaa Xaa Xaa Cys Xaa Cys
 20

```

<210> 12
 <211> 12
 <212> PRT
 <213> Homo sapiens

<400> 12

Leu Gln Asp Pro Cys Ser Asn Cys Pro Ala Gly Thr
1 5 10

Leu Gln Asp Pro Cys Ser Asn Cys Pro Ala Gly Thr
1 5 10